

SEQUENCE LISTING



#10/10

<110> ASHKENAZI, AVI J.  
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 GENENTECH INC.

<120> DcR3 Polypeptide, A TNFR Homolog

<130> 11669.31US03

<140> 09/157,289

<141> 1998-09-18

<150> 60/059,288

<151> 1997-09-18

<150> 60/094,640

<151> 1998-07-30

<160> 16

<170> PatentIn Ver. 2.0

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<211> 300

<212> PRT

<213> Homo sapiens

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Thr Pro Thr Tyr Pro Trp Arg Asp Ala Glu Thr Gly Glu Arg Leu Val  
 35 40 45

Cys Ala Gln Cys Pro Pro Gly Thr Phe Val Gln Arg Pro Cys Arg Arg  
 50 55 60

Asp	Ser	Pro	Thr	Thr	Cys	Gly	Pro	Cys	Pro	Pro	Arg	His	Tyr	Thr	Gln	65	70	75	80
Phe	Trp	Asn	Tyr	Leu	Glu	Arg	Cys	Arg	Tyr	Cys	Asn	Val	Leu	Cys	Gly	85	90	95	
Glu	Arg	Glu	Glu	Glu	Ala	Arg	Ala	Cys	His	Ala	Thr	His	Asn	Arg	Ala	100	105	110	
Cys	Arg	Cys	Arg	Thr	Gly	Phe	Phe	Ala	His	Ala	Gly	Phe	Cys	Leu	Glu	115	120	125	
His	Ala	Ser	Cys	Pro	Pro	Gly	Ala	Gly	Val	Ile	Ala	Pro	Gly	Thr	Pro	130	135	140	
Ser	Gln	Asn	Thr	Gln	Cys	Gln	Pro	Cys	Pro	Pro	Gly	Thr	Phe	Ser	Ala	145	150	155	160
Ser	Ser	Ser	Ser	Ser	Glu	Gln	Cys	Gln	Pro	His	Arg	Asn	Cys	Thr	Ala	165	170	175	
Leu	Gly	Leu	Ala	Leu	Asn	Val	Pro	Gly	Ser	Ser	Ser	His	Asp	Thr	Leu	180	185	190	
Cys	Thr	Ser	Cys	Thr	Gly	Phe	Pro	Leu	Ser	Thr	Arg	Val	Pro	Gly	Ala	195	200	205	
Glu	Glu	Cys	Glu	Arg	Ala	Val	Ile	Asp	Phe	Val	Ala	Phe	Gln	Asp	Ile	210	215	220	
Ser	Ile	Lys	Arg	Leu	Gln	Arg	Leu	Leu	Gln	Ala	Leu	Glu	Ala	Pro	Glu	225	230	235	240
Gly	Trp	Gly	Pro	Thr	Pro	Arg	Ala	Gly	Arg	Ala	Ala	Leu	Gln	Leu	Lys	245	250	255	
Leu	Arg	Arg	Arg	Leu	Thr	Glu	Leu	Leu	Gly	Ala	Gln	Asp	Gly	Ala	Leu	260	265	270	
Leu	Val	Arg	Leu	Leu	Gln	Ala	Leu	Arg	Val	Ala	Arg	Met	Pro	Gly	Leu	275	280	285	
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aggcctgtcg ctgctgtgcc tgggtgttggc gctgcctgcc ctgctgccgg tgcgggctgt 180  
acgcggagtg gcagaaacac ccacctaccc ctggcgggac gcagagacag gggagcggct 240  
ggtgtgcgcc cagtgcctcc caggcacctt tgtgcagcgg ccgtgccgcc gagacagccc 300  
cacgacgtgt ggcccgtgtc caccgcgcc aacacgcagc ttctggaact acctggagcg 360  
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caccacaac cgtgcctgcc gctgcgcgac cggtctcttc gcgcacgctg gtttctgctt 480  
ggagcacgca tcgtgtccac ctggtgccgg cgtgattgcc ccgggcaccc ccagccagaa 540  
cacgcagtgc cagccgtgcc cccagggcac cttctcagcc agcagctcca gctcagagca 600  
gtgccagccc caccgcaact gcacggccct gggcctggcc ctcaatgtgc caggctcttc 660  
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agctgaggag tgtgagcgtg ccgtcatcga ctttgtggct ttccaggaca tctccatcaa 780  
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gcaggacggg gcgctgctgg tgcggctgct gcaggcgctg cgcgtggcca ggatgcccg 960  
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gggcttgcca cgccaccac aaccgtgcct gccgctgccg caccggcttc ttcgcgcacg 180
ctggtttctg cttggagcac gcatcgtgtc cacctgggtgc cggcgtgatt gccccgggca 240
ccccagcca gaacacgcag tgctagccg tgccccccag gcaccttctc agccagcagc 300
tccagctcag agcagtgcc gccccaccgc aactgcacgg ccctgggcct ggccctcaat 360
gtgccaggct cttcctcca tgacaccctg tgcaccagct gcactggctt ccccctcagc 420
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gngcttgcca cgccaccac aaccgcgcct gcngetgcag caccggnttc ttgcgcacg 180
ctgntttctg cttggagcac gcacgtgtgc cacctggtgn cggcgtgatt gcnccgggca 240
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cttcgcgcac gctggtttct gcttgagca cgcacgtgtt ccacctggtg ccggcgtgat 180
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acccccagcc agaacacgca gnccagccgt gccccccagg caccttctca gccagcagct 180
ccagctcaga gcagtgccag cccaccgca actgcacggc cctgggcctg gccctcaatg 240
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cgctgtgcac cagctgcact ggcttcccc tcagcaccag ggtancagga gctgaggagt 180  
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cgcaactgca acgccctggn ctggccctca atgtgccagg ctcttctctc catgacaccc 180
tgtgcaccag ctgcactggc ttccccctca gcaccagggt accaggagct gaggagtgtg 240
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22

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53

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24

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17

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16